<213> homo sapiens

SEQUENCE LISTING

```
<110> Walke, D. Wade
     Mathur, Brian
     Turner, C. Alexander Jr.
     Friddle, Carl Johan
      Gerhardt, Brenda
<120> Novel Human Ion Channel Proteins and Polynucleotides Encoding the
Same
<130> LEX-0208-USA
<150> US 60/221,643
<151> 2000-07-28
<150> US 60/222,503
<151> 2000-08-02
<160> 8
<170> FastSEO for Windows Version 4.0
<210> 1
<211> 1638
<212> DNA
<213> homo sapiens
atgctcaaac agagtgagag gagacggtcc tggagctaca ggccctggaa cacgacggag
                                                                        60
aatgagggca gccaacaccg caggagcatt tgctccctgg gtgcccgttc cggctcccag
                                                                       120
                                                                       180
qccagcatcc acggctggac agagggcaac tataactact acatcgagga agacgaagac
ggsgaggagg aggaccagtg gaaggacgac ctggcagaag aggaccagca ggcaggggag
                                                                       240
                                                                       300
gtcaccaccg ccaagcccga gggccccagc gaccctccgg ccctgctgtc cacgctgaat
gtgaacgtgg gtggccacag ctaccagctg gactactgcg agctggccgg cttccccaag
                                                                       360
acgcgcctag gtcgcctggc cacctccacc agccgcagcc gccagctaag cctgtgcgac
                                                                       420
gactacgagg agcagacaga cgaatacttc ttcgaccgcg acccggccgt cttccagctg
                                                                       480
                                                                       540
gtctacaatt tctacctgtc cggggtgctg ctggtgctcg acgggctgtg tccgcgccgc
ttcctggagg agctgggcta ctggggcgtg cggctcaagt acacgccacg ctgctgccgc
                                                                       600
atctgcttcg aggagcggcg cgacgagctg agcgaacggc tcaagatcca gcacgagctg
                                                                       660
                                                                       720
cgcgcgcagg cgcaggtcga ggaggcggag gaactcttcc gcgacatgcg cttctacggc
ccgcagcggc gccgcctctg gaacctcatg gagaagccrt tctcctcggt ggccgccaag
                                                                       780
                                                                       840
gccatcgggg tggcctccag caccttcgtg ctcgtctccg tggtggcgct ggcgctcaac
                                                                       900
accgtggagg agatgcagca gcactcgggg cagggcgagg gcggcccaga cctgcggccc
                                                                       960
atcctggagc acgtggagat gctgtgcatg ggcttcttca cgctcgagta cctgctgcgc
                                                                      1020
ctagcctcca cgcccgacct gaggcgcttc gcgcgcagcg ccctcaacct ggtggacctg
                                                                      1080
gtggccatcc tgccgctcta ccttcagctg ctgctcgagt gcttcacggg cgagggccac
caacgcggcc agacggtggg cagcgtgggt aaggtgggtc aggtgttgcg cgtcatgcgc
                                                                      1140
                                                                      1200
ctcatgcgca tcttccgcat cctcaagctg gcgcgccact ccaccggact gcgtgccttc
                                                                      1260
ggcttcacgc tgcgccagtg ctaccagcag gtgggctgcc tgctgctctt catcgccatg
                                                                      1320
ggcatcttca ctttctctgc ggctgtctac tctgtggagc acgatgtgcc cagcaccaac
                                                                      1380
ttcactacca tcccccactc ctggtggtgg gccgcggtga gcatctccac cgtgggctac
ggagayatgt acccagagac ccacctgggc aggttttttg ccttcctctg cattgctttt
                                                                      1440
gggatcattc tcaacgggat gcccatttcc atcctctaca acaagttttc tgattactac
                                                                      1500
                                                                      1560
agcaagctga aggcttatga gtataccacc atacgcaggg rgaggggaga ggtgaacttc
atgcagagag ccagaaagaa gatagctgag tgtttgcttg gaagcaaccc acagctcacc
                                                                      1620
ccaagacaag agaattag
                                                                      1638
<210> 2
<211> 545
<212> PRT
```

Met Leu Lys Gln Ser Glu Arg Arg Ser Trp Ser Tyr Arg Pro Trp 1.0 Asn Thr Thr Glu Asn Glu Gly Ser Gln His Arg Arg Ser Ile Cys Ser 2.0 Leu Gly Ala Arg Ser Gly Ser Gln Ala Ser Ile His Gly Trp Thr Glu Gly Asn Tyr Asn Tyr Tyr Ile Glu Glu Asp Glu Asp Gly Glu Glu Glu Asp Gln Trp Lys Asp Asp Leu Ala Glu Glu Asp Gln Gln Ala Gly Glu Val Thr Thr Ala Lys Pro Glu Gly Pro Ser Asp Pro Pro Ala Leu Leu Ser Thr Leu Asn Val Asn Val Gly Gly His Ser Tyr Gln Leu Asp Tyr Cys Glu Leu Ala Gly Phe Pro Lys Thr Arg Leu Gly Arg Leu Ala Thr Ser Thr Ser Arg Ser Arg Gln Leu Ser Leu Cys Asp Asp Tyr Glu Glu Gln Thr Asp Glu Tyr Phe Phe Asp Arg Asp Pro Ala Val Phe Gln Leu Val Tyr Asn Phe Tyr Leu Ser Gly Val Leu Leu Val Leu Asp Gly Leu Cys Pro Arg Arg Phe Leu Glu Glu Leu Gly Tyr Trp Gly Val Arg Leu Lys Tyr Thr Pro Arg Cys Cys Arg Ile Cys Phe Glu Glu Arg Arg Asp Glu Leu Ser Glu Arg Leu Lys Ile Gln His Glu Leu Arg Ala Gln Ala Gln Val Glu Glu Ala Glu Glu Leu Phe Arg Asp Met Arg Phe Tyr Gly Pro Gln Arg Arg Leu Trp Asn Leu Met Glu Lys Pro Phe Ser Ser Val Ala Ala Lys Ala Ile Gly Val Ala Ser Ser Thr Phe Val Leu Val Ser Val Val Ala Leu Ala Leu Asn Thr Val Glu Glu Met Gln Gln His Ser Gly Gln Gly Glu Gly Pro Asp Leu Arg Pro Ile Leu Glu His Val Glu Met Leu Cys Met Gly Phe Phe Thr Leu Glu Tyr Leu Leu Arg Leu Ala Ser Thr Pro Asp Leu Arg Arg Phe Ala Arg Ser Ala Leu Asn Leu Val Asp Leu Val Ala Ile Leu Pro Leu Tyr Leu Gln Leu Leu Leu Glu Cys Phe Thr Gly Glu Gly His Gln Arg Gly Gln Thr Val Gly Ser Val Gly Lys Val Gly Gln Val Leu Arg Val Met Arg Leu Met Arg Ile Phe Arg Ile Leu Lys Leu Ala Arg His Ser Thr Gly Leu Arg Ala Phe Gly Phe Thr Leu Arg Gln Cys Tyr Gln Gln Val Gly Cys Leu Leu Leu Phe Ile Ala Met Gly Ile Phe Thr Phe Ser Ala Ala Val Tyr Ser Val Glu His Asp Val Pro Ser Thr Asn Phe Thr Thr Ile Pro His Ser Trp Trp Trp Ala Ala Val Ser Ile Ser Thr Val Gly Tyr Gly Asp Met Tyr Pro Glu Thr His Leu Gly Arg Phe Phe Ala Phe Leu Cys Ile Ala Phe Gly Ile Ile Leu Asn Gly Met Pro Ile Ser Ile Leu Tyr Asn Lys Phe

```
Ser Asp Tyr Tyr Ser Lys Leu Lys Ala Tyr Glu Tyr Thr Thr Ile Arg
                                505
            500
Arg Glu Arg Gly Glu Val Asn Phe Met Gln Arg Ala Arg Lys Lys Ile
                                                525
        515
                            520
Ala Glu Cys Leu Leu Gly Ser Asn Pro Gln Leu Thr Pro Arg Gln Glu
                                            540
                        535
    530
Asn
545
<210> 3
<211> 180
<212> DNA
<213> homo sapiens
<400> 3
atgctcaaac agagtgagag gagacggtcc tggagctaca ggccctgtcc ggggtgctgc
                                                                        60
                                                                       120
tggtgctcga cgggctgtgt ccgccgct tcctggagga gctgggctac tggggcgtgc
                                                                       180
ggctcaagta cacgccacgc tgctgccgca tctgcttcga ggagcggcgc gacgagctga
<210> 4
<211> 59
<212> PRT
<213> homo sapiens
<400> 4
Met Leu Lys Gln Ser Glu Arg Arg Ser Trp Ser Tyr Arg Pro Cys
                                    10
Pro Gly Cys Cys Trp Cys Ser Thr Gly Cys Val Arg Ala Ala Ser Trp
            20
                                25
Arg Ser Trp Ala Thr Gly Ala Cys Gly Ser Ser Thr Arg His Ala Ala
                            40
Ala Ala Ser Ala Ser Arg Ser Gly Ala Thr Ser
    50
                        55
<210> 5
<211> 2310
<212> DNA
<213> homo sapiens
<400> 5
                                                                        60
tcttcctcta cctcacaggg tcaagggagt gggggaggaa atgggctaag aggttctaaa
                                                                       120
tccctcctaa cacttgcttc ttccaaatca gcaagattag agcagtcaac agctgactgc
gttcagaccc tgcaggctgg gctggcctgc ccaggacctg agaaggggca gctccggtgg
                                                                       180
                                                                       240
caatgtctga gcccctagct gtgctggtcc gggctggcct ctctaagaca gtgcaggcca
cgtgatccat cctcctagag gcagtgagca ggtgagggac ccctacgaca gccaggagga
                                                                       300
                                                                       360
aaaagctagg cgtccacttt ccgcagccat gctcaaacag agtgagagga gacggtcctg
gagctacagg ccctggaaca cgacggagaa tgagggcagc caacaccgca ggagcatttg
                                                                       420
                                                                       480
ctccctgggt gcccgttccg gctcccaggc cagcatccac ggctggacag agggcaacta
                                                                       540
taactactac atcgaggaag acgaagacgg sgaggaggag gaccagtgga aggacgacct
                                                                       600
ggcagaagag gaccagcagg caggggaggt caccaccgcc aagcccgagg gccccagcga
ccctccggcc ctgctgtcca cgctgaatgt gaacgtgggt ggccacagct accagctgga
                                                                       660
ctactgcgag ctggccggct tccccaagac gcgcctaggt cgcctggcca cctccaccag
                                                                       720
                                                                       780
ccgcagccgc cagctaagcc tgtgcgacga ctacgaggag cagacagacg aatacttctt
                                                                       840
cgaccgcgac ccggccgtct tccagctggt ctacaatttc tacctgtccg gggtgctgct
                                                                       900
ggtgctcgac gggctgtgtc cgcgccgctt cctggaggag ctgggctact ggggcgtgcg
gctcaagtac acgccacgct gctgccgcat ctgcttcgag gagcggcgcg acgagctgag
                                                                       960
cgaacggctc aagatccagc acgagctgcg cgcgcaggcg caggtcgagg aggcggagga
                                                                       1020
actettecge gacatgeget tetaeggeee geageggege egeetetgga aceteatgga
                                                                      1080
                                                                      1140
gaagcertte teeteggtgg eegecaagge categgggtg geeteeagea cettegtget
cgtctccgtg gtggcgctgg cgctcaacac cgtggaggag atgcagcagc actcggggca
                                                                      1200
                                                                       1260
gggcgagggc ggcccagacc tgcggcccat cctggagcac gtggagatgc tgtgcatggg
cttcttcacg ctcgagtacc tgctgcgcct agcctccacg cccgacctga ggcgcttcgc
                                                                      1320
gcgcagcgcc ctcaacctgg tggacctggt ggccatcctg ccgctctacc ttcagctgct
                                                                      1380
```

```
1440
gctcgagtgc ttcacgggcg agggccacca acgcggccag acggtgggca gcgtgggtaa
ggtgggtcag gtgttgcgcg tcatgcgcct catgcgcatc ttccgcatcc tcaagctggc
                                                                      1500
                                                                      1560
gcgccactcc accggactgc gtgccttcgg cttcacgctg cgccagtgct accagcaggt
gggctgcctg ctgctcttca tcgccatggg catcttcact ttctctgcgg ctgtctactc
                                                                      1620
tgtggagcac gatgtgccca gcaccaactt cactaccatc ccccactcct ggtggtgggc
                                                                      1680
                                                                      1740
cqcqqtgaqc atctccaccg tgggctacgg agayatgtac ccagagaccc acctgggcag
                                                                      1800
gttttttgcc ttcctctgca ttgcttttgg gatcattctc aacgggatgc ccatttccat
                                                                      1860
cctctacaac aagttttctg attactacag caagctgaag gcttatgagt ataccaccat
acgcagggrg aggggagagg tgaacttcat gcagagagcc agaaagaaga tagctgagtg
                                                                      1920
tttqcttqga agcaacccac agctcacccc aagacaagag aattagtatt ttataggaca
                                                                      1980
                                                                      2040
tgtggctggt agattccatg aacttcaagg cttcattgct ctttttttaa tcattatgat
                                                                      2100
tggcagcaaa aggaaatgtg aagcagacat acacaaaggc catttcgttc acaaagtact
                                                                      2160
qcctctagaa atactcattt tggcccaaac tcagaatgtc tcatagttgc tctgtgttgt
                                                                      2220
gtgaaacatc tgaccttctc aatgacgttg atattgaaaa cctgagggga gcaacagctt
                                                                      2280
agatttttct tgtagcttct cgtggcatct agctcaataa atatttttgg acttgaaaaa
                                                                      2310
aaaaaaaaa aaaaaaaaa aaaaaaaaaa
<210> 6
<211> 1458
<212> DNA
<213> Homo sapiens
<400> 6
                                                                        60
atgagctcag cctgctggga ggccacaggg agatgcaggc tgggcggcgg gtggatggtt
ccaaccggtt gggtccgggg cctggagctc agcctgtggg gtggggaccc agtggtgccc
                                                                       120
tggagctgcc gcttctgctc tcagcaggat gatgggcagg acagggagag gctgacctac
                                                                       180
                                                                       240
ttccagaacc tgcctgagtc tctgacttcc ctcctggtgc tgctgaccac ggccaacaac
                                                                       300
cccgatgtga tgattcctgc gtattccaag aaccgggcct atgccatctt cttcatagtc
ttcactgtga taggaagcct gtttctgatg aacctgctga cagccatcat ctacagtcag
                                                                       360
                                                                       420
ttccggggct acctgatgaa atctctccag acctcgctgt ttcggaggcg gctgggaacc
                                                                       480
cgggctgcct ttgaagtcct atcctccatg gtgggggagg gaggagcctt ccctcaggcc
accegecgag gecegagtac cagteteegt ttetgeagag egeceagtte etetteggee
                                                                       540
                                                                       600
actactatt tgactacctg gggaacctca tcgccctggc aaacctggtg tccatttgcg
                                                                       660
tgttcctggt gctggatgca gatgtgctgc ctgctgagcg tgatgacttc atcctgggga
                                                                       720
tteteaactg egtetteatt gtgtaetaec tgttggagtt getgeteaag gtetttgeec
                                                                       780
tgggcctgcg agggtacctg tcctacccca gcaacgtgtt tgacgggctc ctcaccgttg
tcctgctgga ggccggagat ggtgggcctg ctgtcgctgt gggacatgac ccgcatgctg
                                                                       840
aacatgctca tcgtgttccg cttcctgcgt atcatcccca gcatgaagcc gatggccgtg
                                                                       900
                                                                       960
gtggccagta ccgtcctggg cctggtgcag aacatgcgtg cgtttggcgg gatcctggtg
                                                                      1020
qtqqtctact acqtatttqc catcattggg atcaacttgt ttagaggcgt cattgtggct
                                                                      1080
cttcctggaa acagcagcct ggcccctgcc aatggctcgg cgccctgtgg gagcttcgag
                                                                      1140
cagctggagt actgggccaa caacttcgat gactttgcgg ctgccctggt cactctgtgg
aacttgatgg tggtgaacaa ctggcaggtg tttctggatg catatcggcg ctactcaggc
                                                                      1200
ccgtggtcca agatctattt tgtattgtgg tggctggtgt cgtctgtcat ctgggtcaac
                                                                      1260
ctgtttctgg ccctgattct ggagaacttc cttcacaagt gggacccccg cagccacctg
                                                                      1320
                                                                      1380
cageccettg etgggacece agaggecace taccagatga etgtggaget cetgtteagg
                                                                      1440
gatattctgg aggagcccgg ggaggatgag ctcacagaga ggctgagcca gcacccgcac
                                                                      1458
ctgtggctgt gcaggtga
<210> 7
<211> 485
<212> PRT
<213> Homo sapiens
<400> 7
Met Ser Ser Ala Cys Trp Glu Ala Thr Gly Arg Cys Arg Leu Gly Gly
 1
                                     10
                                                         15
Gly Trp Met Val Pro Thr Gly Trp Val Arg Gly Leu Glu Leu Ser Leu
                                 25
Trp Gly Gly Asp Pro Val Val Pro Trp Ser Cys Arg Phe Cys Ser Gln
        35
                             40
                                                 45
```

Gln Asp Asp Gly Gln Asp Arg Glu Arg Leu Thr Tyr Phe Gln Asn Leu

```
55
Pro Glu Ser Leu Thr Ser Leu Leu Val Leu Leu Thr Thr Ala Asn Asn
                   70
                                        75
Pro Asp Val Met Ile Pro Ala Tyr Ser Lys Asn Arg Ala Tyr Ala Ile
                                    90
               85
Phe Phe Ile Val Phe Thr Val Ile Gly Ser Leu Phe Leu Met Asn Leu
                                105
Leu Thr Ala Ile Ile Tyr Ser Gln Phe Arg Gly Tyr Leu Met Lys Ser
        115
                           120
                                                125
Leu Gln Thr Ser Leu Phe Arg Arg Leu Gly Thr Arg Ala Ala Phe
                       135
                                            140
Glu Val Leu Ser Ser Met Val Gly Glu Gly Gly Ala Phe Pro Gln Ala
                    150
                                        155
Thr Arg Arg Gly Pro Ser Thr Ser Leu Arg Phe Cys Arg Ala Pro Ser
               165
                                    170
                                                        175
Ser Ser Ser Ala Thr Thr Thr Leu Thr Thr Trp Gly Thr Ser Ser Pro
           180
                               185
                                                   190
Trp Gln Thr Trp Cys Pro Phe Ala Cys Ser Trp Cys Trp Met Gln Met
                           200
                                                205
Cys Cys Leu Leu Ser Val Met Thr Ser Ser Trp Gly Phe Ser Thr Ala
                       215
                                            220
Ser Ser Leu Cys Thr Thr Cys Trp Ser Cys Cys Ser Arg Ser Leu Pro
                    230
                                        235
Trp Ala Cys Glu Gly Thr Cys Pro Thr Pro Ala Thr Cys Leu Thr Gly
                245
                                    250
Ser Ser Pro Leu Ser Cys Trp Arg Pro Glu Met Val Gly Leu Leu Ser
           260
                                265
                                                    270
Leu Trp Asp Met Thr Arg Met Leu Asn Met Leu Ile Val Phe Arg Phe
                            280
                                                285
Leu Arg Ile Ile Pro Ser Met Lys Pro Met Ala Val Val Ala Ser Thr
                        295
                                            300
Val Leu Gly Leu Val Gln Asn Met Arg Ala Phe Gly Gly Ile Leu Val
                    310
                                        315
Val Val Tyr Tyr Val Phe Ala Ile Ile Gly Ile Asn Leu Phe Arg Gly
                325
                                    330
                                                        335
Val Ile Val Ala Leu Pro Gly Asn Ser Ser Leu Ala Pro Ala Asn Gly
           340
                                345
                                                    350
Ser Ala Pro Cys Gly Ser Phe Glu Gln Leu Glu Tyr Trp Ala Asn Asn
     355
                           360
                                               365
Phe Asp Asp Phe Ala Ala Ala Leu Val Thr Leu Trp Asn Leu Met Val
                       375
                                            380
Val Asn Asn Trp Gln Val Phe Leu Asp Ala Tyr Arg Arg Tyr Ser Gly
                   390
                                       395
                                                           400
Pro Trp Ser Lys Ile Tyr Phe Val Leu Trp Trp Leu Val Ser Ser Val
                                    410
Ile Trp Val Asn Leu Phe Leu Ala Leu Ile Leu Glu Asn Phe Leu His
                                425
Lys Trp Asp Pro Arg Ser His Leu Gln Pro Leu Ala Gly Thr Pro Glu
        435
                            440
                                                445
Ala Thr Tyr Gln Met Thr Val Glu Leu Leu Phe Arg Asp Ile Leu Glu
                       455
                                           460
Glu Pro Gly Glu Asp Glu Leu Thr Glu Arg Leu Ser Gln His Pro His
                                        475
Leu Trp Leu Cys Arg
```

<210> 8

<211> 2905

<212> DNA

<213> Homo sapiens

<400> 8

gagtcggctg	cgagcaggcg	aggtggcctg	agggaggtca	ctaggctggc	tgagggcttt	120
ttgctgtggt	tctgagccgg	cctgcttcca	ggcaccgtgt	ccatgcgggt	gagcggtctc	180
cctgggtgcc	cactcttgcg	cccggagatc	ctgagtttgg	tcctgtctgg	ccatgagctc	240
agcctgctgg	gaggccacag	ggagatgcag	gctgggcggc	gggtggatgg	ttccaaccgg	300
ttgggtccgg	ggcctggagc	tcagcctgtg	gggtggggac	ccagtggtgc	cctggagctg	360
ccgcttctgc	tctcagcagg	atgatgggca	ggacagggag	aggctgacct	acttccagaa	420
cctgcctgag	tctctgactt	ccctcctggt	gctgctgacc	acggccaaca	accccgatgt	480
gatgattcct	gcgtattcca	agaaccgggc	ctatgccatc	ttcttcatag	tcttcactgt	540
gataggaagc	ctgtttctga	tgaacctgct	gacagccatc	atctacagtc	agttccgggg	600
ctacctgatg	aaatctctcc	agacctcgct	gtttcggagg	cggctgggaa	cccgggctgc	660
ctttgaagtc	ctatcctcca	tggtgggga	gggaggagcc	ttccctcagg	ccacccgccg	720
aggcccgagt	accagtctcc	gtttctgcag	agcgcccagt	tcctcttcgg	ccactactac	780
tttgactacc	tggggaacct	catcgccctg	gcaaacctgg	tgtccatttg	cgtgttcctg	840
gtgctggatg	cagatgtgct	gcctgctgag	cgtgatgact	tcatcctggg	gattctcaac	900
tgcgtcttca	ttgtgtacta	cctgttggag	ttgctgctca	aggtctttgc	cctgggcctg	960
cgagggtacc	tgtcctaccc	cagcaacgtg	tttgacgggc	tcctcaccgt	tgtcctgctg	1020
gaggccggag	atggtgggcc	tgctgtcgct	gtgggacatg	acccgcatgc	tgaacatgct	1080
	cgcttcctgc					1140
	ggcctggtgc					1200
	gccatcattg					1260
	ctggcccctg					1320
	aacaacttcg					1380
	aactggcagg					1440
caagatctat	tttgtattgt	aataactaat	atcatctatc	atctgggtca	acctgtttct	1500
	ctggagaact					1560
	ccagaggcca					1620
adaddadccc	ggggaggatg	agctcacaga	gaggetgage	cagcacccgc	acctgtggct	1680
	cgtccgggct					1740
	catcatggaa					1800
ttcctctgac	ggaccactaa	gctggggaca	ggaaccaagt	cctttacata	tggcccaaca	1860
	gaacagctgc					1920
	agtgagaatt					1980
	gcctcccctg					2040
	caggcctgac					2100
	tgctggtggt					2160
	tccgatttta					2220
	aggccctttc					2280
	atgtccccag					2340
	tggggctttg					2400
	tcagcaaaca					2460
	aaaatctaca					2520
	aaaaacagca					2580
	gccaaaatta					2640
	atcatgaggt					2700
	aaaatacaaa					2760
tactggggag	gctgaggcag	gagaatggcg	tgaacccggg	aagcggagct.	tacaataaac	2820
	ccactgcagt					2880
	aaaaaaaaaa		2355090			2905
aaaaaaaaaaa						